SEQUENCE LISTING

<110> Reed, John C.
Okada, Kazuya

<120> Survivin-Binding Proteins, Encoding Nucleic Acids, and Methods of Use

<130> P-LJ 5144

<150> US 09/770,219
<151> 2001-01-25

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gtcacagaca agacagcaag cagg atg gag cac tac cgg aaa gct ggc tct 171
Met Glu His Tyr Arg Lys Ala Gly Ser
1 5

gta gag ctc cca gcg cct tcc cca atg ccc cag cta cct cct gat acc 219
Val Glu Leu Pro Ala Pro Ser Pro Met Pro Gln Leu Pro Pro Asp Thr
10 15 20 25

ctt gag atg cgg gtc cga gat ggc agc aaa att cgc aac ctg ctg ggg 267
Leu Glu Met Arg Val Arg Asp Gly Ser Lys Ile Arg Asn Leu Leu Gly
30 35 40

ttg gct ctg ggt cgg ttg gag ggc ggc agt gct cgg cat gta gtg ttc 315
Leu Ala Leu Gly Arg Leu Glu Gly Gly Ser Ala Arg His Val Val Phe
45 50 55

tca ggt tct ggc agg gct gca gga aag gct gtc agc tgc gct gag att 363
Ser Gly Ser Gly Arg Ala Ala Gly Lys Ala Val Ser Cys Ala Glu Ile
60 65 70

gtc aag cgg cgg gtc cca ggc ctg cac cag ctc acc aag cta cgt ttc 411
Val Lys Arg Arg Val Pro Gly Leu His Gln Leu Thr Lys Leu Arg Phe

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     75                  80                  85

ctt cag act gag gac agc tgg gtc cca gcc tca cct gac aca ggg cta   459
Leu Gln Thr Glu Asp Ser Trp Val Pro Ala Ser Pro Asp Thr Gly Leu
 90                  95                 100                 105

gac ccc ctc aca gtg cgc cgc cat gtg cct gca gtg tgg gtg ctg ctc   507
Asp Pro Leu Thr Val Arg Arg His Val Pro Ala Val Trp Val Leu Leu
                110                 115                 120

agc cgg gac ccc ctg gac ccc aat gag tgt ggt tac caa ccc cca gga   555
Ser Arg Asp Pro Leu Asp Pro Asn Glu Cys Gly Tyr Gln Pro Pro Gly
            125                 130                 135

gca ccc cct ggc ctg ggt tcc atg ccc agc tcc agc tgt ggc cct cgt   603
Ala Pro Pro Gly Leu Gly Ser Met Pro Ser Ser Ser Cys Gly Pro Arg
        140                 145                 150

tcc cga aga agg ctc gag aca ccc gat cgt gaa gac ttg tga           645
Ser Arg Arg Arg Leu Glu Thr Pro Asp Arg Glu Asp Leu
    155                 160                 165
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<213> Homo sapiens

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Gly Ser Lys Ile Arg Asn Leu Leu Gly Leu Ala Leu Gly Arg Leu Glu
        35                  40                  45
Gly Gly Ser Ala Arg His Val Val Phe Ser Gly Ser Gly Arg Ala Ala
    50                  55                  60
Gly Lys Ala Val Ser Cys Ala Glu Ile Val Lys Arg Arg Val Pro Gly
65                  70                  75                  80
Leu His Gln Leu Thr Lys Leu Arg Phe Leu Gln Thr Glu Asp Ser Trp
                85                  90                  95
Val Pro Ala Ser Pro Asp Thr Gly Leu Asp Pro Leu Thr Val Arg Arg
            100                 105                 110
His Val Pro Ala Val Trp Val Leu Leu Ser Arg Asp Pro Leu Asp Pro
        115                 120                 125
Asn Glu Cys Gly Tyr Gln Pro Pro Gly Ala Pro Pro Gly Leu Gly Ser
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Pro Asp Arg Glu Asp Leu
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Pro His Ile Leu Leu Phe Arg Arg Pro
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<211> 41
<212> PRT
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Val Pro Lys Thr His Leu Met Thr Glu Ala Glu Trp Arg Ser Ile Gly
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Pro His Ile Leu Leu Phe Arg Arg Pro
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Pro His Ile Leu Leu Phe Lys Arg Pro

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cca atg ccc cag cta cct cct gat acc ctt gag atg cgg gtc cga gat      96
Pro Met Pro Gln Leu Pro Pro Asp Thr Leu Glu Met Arg Val Arg Asp
            20                  25                  30

ggc agc aaa att cgc aac ctg ctg ggg ttg gct ctg ggt cgg ttg gag     144
Gly Ser Lys Ile Arg Asn Leu Leu Gly Leu Ala Leu Gly Arg Leu Glu
        35                  40                  45

ggc ggc agt gct cgg cat gta gtg ttc tca ggt tct ggc agg gct gca     192
Gly Gly Ser Ala Arg His Val Val Phe Ser Gly Ser Gly Arg Ala Ala
    50                  55                  60

gga aag gct gtc agc tgc gct gag att gtc aag cgg cgg gtc cca ggc     240
Gly Lys Ala Val Ser Cys Ala Glu Ile Val Lys Arg Arg Val Pro Gly
65                  70                  75                  80

ctg cac cag ctc acc aag cta cgt ttc ctt cag act gag gac agc tgg     288
Leu His Gln Leu Thr Lys Leu Arg Phe Leu Gln Thr Glu Asp Ser Trp
                85                  90                  95

gtc cca gcc tca cct gac aca ggg cta gac ccc ctc aca gtg cgc cgc     336
Val Pro Ala Ser Pro Asp Thr Gly Leu Asp Pro Leu Thr Val Arg Arg
            100                 105                 110
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cat gtg cct gca gtg tgg gtg ctg ctc agc cgg gac ccc ctg gac ccc 384
His Val Pro Ala Val Trp Val Leu Leu Ser Arg Asp Pro Leu Asp Pro
115 120 125

aat gag tgt ggt tac caa ccc cca gga gca ccc cct ggc ctg ggt tcc 432
Asn Glu Cys Gly Tyr Gln Pro Pro Gly Ala Pro Pro Gly Leu Gly Ser
130 135 140

atg ccc agc tcc agc tgt ggc cct cgt tcc cga aga agg gct cga gac 480
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acc cga tcg tga 492
Thr Arg Ser

<210> 14
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<213> Homo sapiens

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Gly Ser Lys Ile Arg Asn Leu Leu Gly Leu Ala Leu Gly Arg Leu Glu
35 40 45
Gly Gly Ser Ala Arg His Val Val Phe Ser Gly Ser Gly Arg Ala Ala
50 55 60
Gly Lys Ala Val Ser Cys Ala Glu Ile Val Lys Arg Arg Val Pro Gly
65 70 75 80
Leu His Gln Leu Thr Lys Leu Arg Phe Leu Gln Thr Glu Asp Ser Trp
85 90 95
Val Pro Ala Ser Pro Asp Thr Gly Leu Asp Pro Leu Thr Val Arg Arg
100 105 110
His Val Pro Ala Val Trp Val Leu Leu Ser Arg Asp Pro Leu Asp Pro
115 120 125
Asn Glu Cys Gly Tyr Gln Pro Pro Gly Ala Pro Pro Gly Leu Gly Ser
130 135 140
Met Pro Ser Ser Ser Cys Gly Pro Arg Ser Arg Arg Arg Ala Arg Asp
145 150 155 160
Thr Arg Ser